

Replacement Sheet

Figure 15A

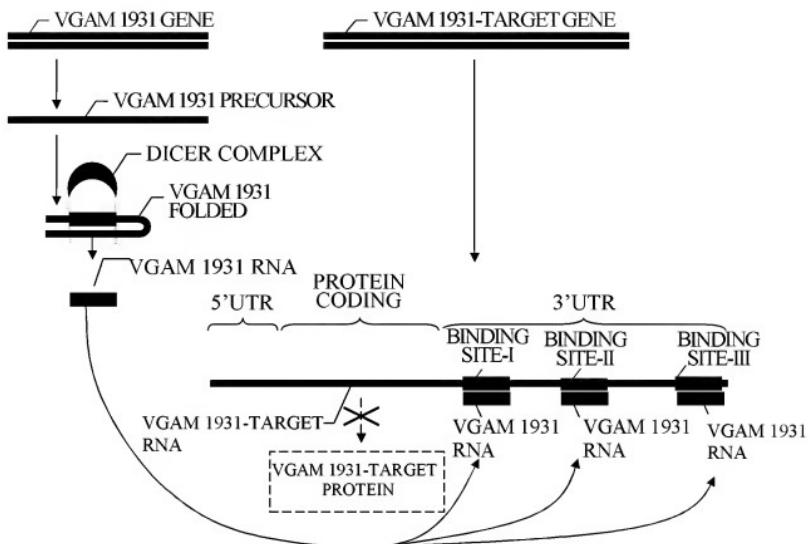


Figure 15B

CCCGTGGGGGTCTTAGTGAAGTGACGTGCTGTGAATA **SEQ ID:1917**
CAGGTCCATAGCACCGCTATCCACTATGTCTGCCCGGG

TAGCACCGCTATCCACTATGTCTC

SEQ ID: 4642

Figure 15C

T	G	TCT	-	AC	A	A
CCC	GG	GGG	TAGTGG	AGTG	GTGCTG	AT C
GGGC	CC	CTC	ATCACCT	<u>TCGC</u>	CACGATAC	TG A
-	G	<u>TGT</u>	A	--	C	G

Replacement Sheet

Figure 15D/1

	CT-	CACTAT		
COL6A1 BINDING SITE1	5' TAGCACCG 3'	ATC TAG AAT	GTCT CAGA AGCTCC	3' 5' SEQ ID: 4642 SEQ ID: 7584
	CC	AT-		
SFRS1 BINDING SITE1	5' GCA 3'	GCTATCCACT CGT CGATGGGTGG AA	GTC CAG ATT	3' 5' SEQ ID: 4642 SEQ ID: 13801
	A-	CTAT	A	
HIP12 BINDING SITE1	5' TAGC 3'	CCG GGC CG	CC GG CC--	3' 5' SEQ ID: 4642 SEQ ID: 32922
	CTA	ACTA	-	
ZNF212 BINDING SITE1	5' TAGCACCG 3'	TCC AGG AC-	TG AC GG--	3' 5' SEQ ID: 4642 SEQ ID: 14557
	A	C--	A	
FLJ20436 BINDING SITE1	5' GC 3'	CCG GGT - TTT	TATCCACT ATAGGTGA C	3' 5' SEQ ID: 4642 SEQ ID: 19472
	C -	A	CAC	
KIAA1622 BINDING SITE1	5' TAGCA 3'	C GCT TC ATTGT G CGA AG A T C TA-	TATGTCTC ATACAGAG	3' 5' SEQ ID: 4642 SEQ ID: 27766
	ACC	C	G	
LOC51312 BINDING SITE1	5' GC 3'	GCTATCCA CGATAGGT A--	TAT TC ATA AG - A	3' 5' SEQ ID: 4642 SEQ ID: 20659
	AC	C	TCCA	
LOC57105 BINDING SITE1	5' TAGC 3'	CG TA ATCG GC AT CA A	CTATGTCTC GATATAGAG CAAC	3' 5' SEQ ID: 4642 SEQ ID: 21639
	CC	A		
LOC146603 BINDING SITE1	5' GCA 3'	GCT TGT CGA TA	TCCACTATGT AGGTGATACA C	3' 5' SEQ ID: 4642 SEQ ID: 38215

Replacement Sheet

Figure 15D/2

	G ATC	A			
LOC145761	5' ACC CT	CACT	TGTCTC	3'	SEQ ID:4642
BINDING SITE1					
	3' TGG GA	GTGA	ACAGAG	5'	SEQ ID:40584
	G AAA	C			
	---	CT	CAC		
LOC202986	5' TAGCA CCG	ATC	TATGTCTC	3'	SEQ ID:4642
BINDING SITE1					
	3' ATCGT GGT	TAG	ATACAGAG	5'	SEQ ID:43470
	TA	T-	CA-		